

Notice of References Cited	Application/Control No. 10/791,017	Applicant(s)/Patent Under Reexamination OBENDORF ET AL.	
	Examiner Zachary C. Howard	Art Unit 1646	Page 1 of 2

U.S. PATENT DOCUMENTS

*		Document Number Country Code-Number-Kind Code	Date MM-YYYY	Name	Classification
*	A	US-2003/0082511 A1	05-2003	Brown et al.	435/4
	B	US-			
	C	US-			
	D	US-			
	E	US-			
	F	US-			
	G	US-			
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	I	US-			
	J	US-			
	K	US-			
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FOREIGN PATENT DOCUMENTS

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NON-PATENT DOCUMENTS

*		Include as applicable: Author, Title Date, Publisher, Edition or Volume, Pertinent Pages)
	U	Wells (18 September 1990) "Additivity of Mutational Effects in Proteins." Biochemistry 29(37): 8509-8517. ✓
	V	Ngo et al. (2 March 1995) "The Protein Folding Problem and Tertiary Structure Prediction, Chapter 14: Computational Complexity Protein Structure Prediction, and the Levinthal Paradox" pp. 492-495. ✓
	W	Bork (2000) "Powers and Pitfalls in Sequence Analysis: The 70% Hurdle." Genome Research 10:398-400. ✓
	X	Skolnick and Fetrow (2000) "From gene to protein structure and function: novel applications of computational approaches in the genomic era." Trends in Biotech. 18(1): 34-39. ✓

*A copy of this reference is not being furnished with this Office action. (See MPEP § 707.05(a).)
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	U	Doerks et al. (June 1998) "Protein annotation: detective work for function prediction." Trends in Genetics 14(6): 248-250. ✓
	V	Smith and Zhang (November 1997) "The challenges of genome sequence annotation or 'The devil is in the details'." Nature Biotechnology 15:1222-1223. ✓
	W	Brenner (April 1999) "Errors in genome annotation." Trends in Genetics 15(4): 132-133. ✓
	X	Bork and Bairoch (October 1996) "Go hunting in sequence databases but watch out for the traps." Trends in Genetics 12(10): 425-427. ✓

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